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# PBL Project 2

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% Code Contributors:
% Anika Kulkarni      Last Commit: 10/21/2025
% Krishna Ravi       Last Commit: 10/20/2025
% Felix Wang         Last Commit: 10/21/2025

% Overview
% -----
% ACCOUNTING EQUATION:  $\Psi_{out} = \Psi_{in} [1 - w_{reabs} + w_{sec}]$ 

% ASSUMPTIONS:
% (1) Steady-State
% (2) No Reactions
% (3) No Leaks

% LIMITATIONS:
% (1) Water flow held constant (no ADH modulation)
% (2) Active/passive Transport not explicitly modeled
% (3) Tubuloglomerular feedback not included

% OUTPUTS:
% > Model allows for various scenarios
% (1) Mass flow rate of each chemical constituent in each unit
% (2) Concentration of each chemical constituent in each unit

% Kidney Model
% -----
function[molar_flow_rates, concs, grams_per_min_out] = kidney_model(C0,
snGFR, condition)

% Indices
% -----
% Nephronal Units (x-Axis)
units = ["RC", "PT(S1)", "PT(S2)", "PT(S3)", "DL", "AL", "DT", "CD"];
% RC = Renal Corpuscle
% PT = Proximal Tubule: Split into S1, S2, S3
% DL = Descending Limb
% AL = Ascending Limb
% DT = Distal Tubule
% CD = Collecting Duct

% Chemical Constituents (y-Axis)
chemicals = ["Na^+", "Cl^-", "Urea", "Glucose", "K^+", "HCO_3^-", "Mg^{2+}",
"PO_4^{3-}", "Creatinine", "Ca^{2+}"];
colors = ["#012966", "#005f73", "#0a9396", "#94d2bd", "#e9d8a6", "#ee9b00",
"#ca6702", "#bb3e03", "#ae2012", "#9b2226"];
molec_weights = [22.989, 35.453, 60.056, 180.156, 39.098, 61.020, 24.305,
```

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```
94.971, 113.12, 40.08]; % g/mol
```

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% Matrix Lengths
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nSeg = length(units);
```

```
nSol = length(chemicals);
```

```
% Input Data Matrices
```

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% -----
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```
% Fractions derived from a host of literature sources, primarily:
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```
% - Weinstein, A. M. Seldin and Giebisch's The Kidney. Elsevier Inc., 2008.  
849-887.1081-1142
```

```
% - Vallon, V. Am J Physiol Cell Physiol. 2011 Jan; 300(1): C6-C8.
```

```
% - Oregon State University, open courseware
```

```
%{
```

```
      Na+      Cl-      Urea      Gluc      K+      HCO3-      Mg2+      PO4^3-  
Creat      Ca2+  
RC | (1,1) | (1,2) | (1,3) | (1,4) | (1,5) | (1,6) | (1,7) | (1,8) | (1,9) |  
(1,10)  
S1 | (2,1) | (2,2) | (2,3) | (2,4) | (2,5) | (2,6) | (2,7) | (2,8) | (2,9) |  
(2,10)  
S2 | (3,1) | (3,2) | (3,3) | (3,4) | (3,5) | (3,6) | (3,7) | (3,8) | (3,9) |  
(3,10)  
S3 | (4,1) | (4,2) | (4,3) | (4,4) | (4,5) | (4,6) | (4,7) | (4,8) | (4,9) |  
(4,10)  
DL | (5,1) | (5,2) | (5,3) | (5,4) | (5,5) | (5,6) | (5,7) | (5,8) | (5,9) |  
(5,10)  
AL | (6,1) | (6,2) | (6,3) | (6,4) | (6,5) | (6,6) | (6,7) | (6,8) | (6,9) |  
(6,10)  
DT | (7,1) | (7,2) | (7,3) | (7,4) | (7,5) | (7,6) | (7,7) | (7,8) | (7,9) |  
(7,10)  
CD | (8,1) | (8,2) | (8,3) | (8,4) | (8,5) | (8,6) | (8,7) | (8,8) | (8,9) |  
(8,10)  
%}
```

```
% Reabsorption Fractions per Constituent (of incoming stream) per Unit
```

```
% This is what's returned to the interstitial fluid
```

```
% Solutes      Na+,      Cl-,      Urea,      Glucose, K+,      HCO3-,      Mg2+,  
PO4^3-, Creat, Ca2+  
reabs_frac = [0,      0,      0,      0,      0,      0,      0,  
0,      0,      0; % RC  
0.35,      0,      0.33,      0.33,      0,      0.90,      0,      0.80,      0,  
0.35; % PT(S1)  
0.328,      0.328,      0,      0.09,      0,      0.10,      0,  
0.25; % PT(S2)  
0.222,      0.222,      0.50,      0,      0.60,      0,      0.20,  
0.10; % PT(S3)  
0,      0,      0,      0,      0,      0.15,      0,  
0,      0,      0; % DL  
0.25,      0.25,      0,      0,      0.25,      0,      0.7,  
0,      0,      0.20; % AL  
0.05,      0.05,      0,      0,      0,      0.05,      0.05,  
0.05,      0,      0.10; % DT
```

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0,      0,      0,      0,      0,      0,      0,      0,
0,      0,      0.05; % CD
    ];
%{
    • Note: Na/Cl reabsorption can increase to 0.02/0.03 in the Collecting Duct
when producing dilute urine (over-hydration) or with high salt intake (need
to dilute urine)
    • Note: Urea is reabsorbed when producing concentrated urine to conserve
water, as urea establishes an osmotic gradient (dehydration)
    • However, because urea reabsorption is tightly regulated by ADH and other
hormones, this is outside the scope of this model
%}

% Secretion Fractions per Constituent (of incoming stream) per Unit
% This is what's secreted by the bloodstream into tubules post-RC (initial
filtration)
% Solutes      Na+,      Cl-,      Urea,      Glucose, K+,      HCO3-,      Mg2+,
PO4^3-, Creat,  Ca2+
sec_frac = [0,      0,      0,      0,      0,      0,      0,
0,      0,      0; % RC
            0,      0,      0,      0,      0,      0,      0,
0,      0,      0; % PT(S1)
            0,      0,      0,      0,      0,      0,      0,
0,      0.30,  0; % PT(S2)
            0,      0,      0,      0,      0,      0,      0,
0,      0,      0; % PT(S3)
            0,      0,      0.15,  0,      0,      0,      0,
0,      0,      0; % DL
            0,      0,      0,      0,      0,      0,      0,
0,      0,      0; % AL
            0,      0,      0,      0,      0.10,  0,      0,
0,      0,      0; % DT
            0,      0,      0,      0,      0.10,  0,      0,
0,      0,      0; % CD
    ];

% Safety Clamps
reabs_frac = max(0, min(reabs_frac, 0.999));
sec_frac = max(0, min(sec_frac, 0.999));

% Calculated Data Matrices
% -----
% Molar Flow Rates per Constituent per Unit
molar_flow_rates = zeros(nSeg,nSol); % mmol/min

% Mass Flow Rates per Constituent leaving Collecting Duct (Outlet)
grams_per_min_out = zeros(1,nSol); % g/min

% Concentrations per Constituent per Unit
concs = zeros(nSeg,nSol); % mmol/L = mM

% Volumetric flow rates per constituent per unit
vol_flow_rates = snGFR * ones(nSeg,1); % mL/min

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% Input Parameters
% -----
% Initializes input stream (C0 & snGFR) to Renal Corpuscle
%{
    molar_flow_rates = vol_flow_rates * concs
    mmol/min          = mL/min * mM * 1L/1000mL
%}

concs(1,:) = C0; % mmol/L = mM
molar_flow_rates(1,:) = snGFR .* concs(1,:) * 1e-3; % mmol/min

% Adjustments
% -----
% Adjusts secretion fractions in the Collecting Duct for Na+ in the event of
high salt intake
if concs(1,1) > 140 && concs(1,2) > 106
    sec_frac(8,1) = 0.02; % Na+
    sec_frac(8,2) = 0.03; % Cl-
end

% Calculations
% -----
% Calculates molar flow rates and concentrations for each unit
for i = 2:nSeg
    % Accounting for Each Constituent
    N_in = molar_flow_rates(i-1,:); % Vector with inflows for all
constituents
    N_out = N_in .* (1 - reabs_frac(i,:) + sec_frac(i,:)); % Vector with
outflows for all constituents

    if any(N_out < 0)
        errordlg("Negative Flow Rate at Row " + i + "!!!") % Error Warning
    end

    % Populates row i of Data Matrix
    molar_flow_rates(i,:) = N_out; % mmol/min
    concs(i,:) = molar_flow_rates(i,:) ./ vol_flow_rates(i); % mmol/min /
mL/min = mmol/L
end

% Converts outlet molar flow rates to mass flow rates
for j = 1:nSol
    grams_per_min_out(j) = molar_flow_rates(end,j) * molec_weights(j) *
1e-3; % mmol/min * g/mol * 1e-3 = g/min
end

% Print
% -----
% Molar Flow Rates

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fprintf("\n\n" + condition + "\n-----\n");

fprintf("8x10 Molar Flow Rate Matrix\n" + ...
    "• Rows = RC, S1, S2, S3, DL, AL, DT, CD\n" + ...
    "• Columns = Na+, Cl-, Urea, Glucose, K+, HCO3-, Mg2+, PO4^3-,
Creatinine, Ca2+");
disp(molar_flow_rates);

% Concentrations
fprintf("8x10 Concentrations Matrix\n" + ...
    "• Rows = RC, S1, S2, S3, DL, AL, DT, CD\n" + ...
    "• Columns = Na+, Cl-, Urea, Glucose, K+, HCO3-, Mg2+, PO4^3-,
Creatinine, Ca2+");
disp(concs);

% Outlet Mass Flow Rates
disp("Outlet (collecting duct) in grams/min per solute: ");
for k = 1:nSol
    disp(chemicals(k) + ": " + num2str(grams_per_min_out(k)));
end

% Graphs
% -----
fig = figure('Name', condition, 'Units', 'normalized', 'Position', [0.05
0.05 0.9 0.8]);

% Adds tabs to figure
tabGroup = uitabgroup(fig);
tab1 = uitab(tabGroup, 'Title', 'Concentrations');
tab2 = uitab(tabGroup, 'Title', 'Molar Flow Rates');
tab3 = uitab(tabGroup, 'Title', 'Outlet Mass Flow Rates');

% Display layout for graphs
tLayout1 = tiledlayout(tab1, 2, 5, 'TileSpacing', 'compact', 'Padding',
'compact');
tLayout2 = tiledlayout(tab2, 2, 5, 'TileSpacing', 'compact', 'Padding',
'compact');

% Figure 1: Concentrations
for k = 1:nSol
    ax1 = nexttile(tLayout1);
    plot(ax1, 1:nSeg, concs(:,k), '-', 'Color', colors(k), 'LineWidth', 2);
    grid on
    title(chemicals(k) + " Concentration", 'FontWeight','bold', 'FontSize',
12)
    xticks(1:nSeg)
    xticklabels(units)
    ylabel('mM')
    ylim([0 inf])
end
sgtitle({"Solute Concentrations Along Nephron Segments", "Test Case: " +
condition}, 'FontSize', 14, 'FontWeight','bold')

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% Figure 2: Molar Flow Rates
for k = 1:nSol
    ax2 = nexttile(tLayout2);
    plot(ax2, 1:nSeg, molar_flow_rates(:,k), '-', 'Color', colors(k),
'LineWidth', 2);
    grid on
    title(chemicals(k) + " Molar Flow", 'FontWeight','bold', 'FontSize', 12)
    xticks(1:nSeg)
    xticklabels(units)
    ylabel('mmol/min')
    ylim([0 inf])
end
sgtitle({"Molar Flow Rates Along Nephron Segments", "Test Case: " +
condition}, 'FontSize', 14, 'FontWeight','bold')

% Figure 3: Outlet Mass Flow Rates
axes('Parent', tab3);
b = bar(1:nSol, grams_per_min_out);
b.FaceColor = 'flat';
for k = 1:nSol
    b.CData(k,:) = hex2rgb(colors(k));
end
grid on
xticks(1:nSol)
xticklabels(chemicals)
ylabel('g/min')
ylim([0 inf])
sgtitle({"Outlet Mass Flow Rates Along Nephron Segments", "Test Case: " +
condition}, 'FontSize', 14, 'FontWeight','bold')

end

% Test Cases
% -----
function[conc_out, snGFR_out] = test_cases(C0, snGFR, condition)
% Implements test cases by adjusting inlet concentrations and single-nephron
GFR accordingly
% Test Cases: 'healthy',
%             'ckd3b' (CKD, stage 3b),
%             't2dm_early' (early type 2 diabetes),
%             't2dm_late' (late type 2 diabetes),
%             'htn' (hypertension)

% Re-indexing for convenience
Na = 1;
Cl = 2;
Urea = 3;
Gluc = 4;
K = 5;
HCO3 = 6;
Mg = 7;
PO4 = 8;
Creat = 9;
Ca = 10;

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switch lower(condition)
    case 'healthy'
        % No change, using base values

    case 'ckd3b'
        % We are modeling stage 3b as this marks substantial loss of kidney
function and sure-fire diagnosis of CKD

        % Normal GFR = 90-120 mL/min; Stage 3b GFR = 30-44 mL/min
        % Scaling factor as reference as snGFR values not available
        snGFR = snGFR * 37/105; % Using a scaling factor as reference as
snGFR values not available

        % Creatinine level increased by 60%
        % Typical creatinine level: 0.7-1.3 mg/dL in males, 0.6-1.1 mg/dL in
females
        % CKD3b creatinine level: 1.2-2.0 mg/dL in males, 1.8-3.0 mg/dL in
females
        % 50-70% increase
        C0(Creat) = C0(Creat) * 1.6;

        % Urea level increased by 40% (estimate)
        C0(Urea) = C0(Urea) * 1.4;

        % Potassium level increased by 15% (estimate; hyperkalemia)
        C0(K) = C0(K) * 1.15;

        % Phosphate level increased by 25% (estimate; due to degradation of
phosphate secretion)
        C0(PO4) = C0(PO4) * 1.25;

        % Bicarbonate level decreased, not significantly (sign of metabolic
acidosis)
        C0(HCO3) = C0(HCO3) * 0.90;

        % Calcium level decreased, not significantly (sign of hypocalcemia)
        C0(Ca) = C0(Ca) * 0.90;

    case 't2dm_early'
        % Separating early and late stages because early T2DM =
hyperfiltration

        % Normal GFR = 120 mL/min; Early T2DM GFR = 120-150 mL/min, median
at 135 mL/min
        % Approximating hyperfiltration
        snGFR = snGFR * 1.125;

        % Average plasma concentration of glucose during early hyperglycemia
        C0(Gluc) = 7; % 7 mmol/L = 126 mg/dL

    case 't2dm_late'
        % Separating early and late stages because late T2DM = hypofiltration

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        % Normal GFR = 90-120 mL/min; Late T2DM GFR = 15-29 mL/min, median
at 22 mL/min (correlated with Stage 4-5 CKD)
        % Approximating hypofiltration
        snGFR = snGFR * 22/105;

        % Average plasma concentration of glucose during late stage
hyperglycemia
        C0(Gluc) = 9.7; % 8.3-11.1 mmol/L

        % Potassium level jumps to greater than 6 mmol/L
        C0(K) = C0(K) * 1.30; % Jumps to greater than 6 mmol/L

        % Phosphate level jumps to greater than 1.5 mmol/L
        C0(PO4) = C0(PO4) * 1.15; % Jumps to greater than 1.5 mmol/L

        % Bicarbonate level decreases but still indicative of metabolic
acidosis
        C0(HCO3) = C0(HCO3) * 0.85;

        case 'htn'
            % Slight decrease to reflect vascular damage
            snGFR = snGFR * 0.90;
        end

% Outputs
conc_out = C0;
snGFR_out = snGFR;

end

% Main Code
% -----
% BASELINE INLET CONCENTRATIONS
% Inlet filtrate concentrations at renal corpuscle, as INPUT into main
function

% Na+, Cl-, Urea, Glucose, K+, HCO3-, Mg2+, PO4^3-, Creatinine, Ca2+
C0 = [140.0000, 102.0000, 5.714, 4.6905, 4.3500, 24.0000, 0.8225, 0.3950,
0.0920, 0.5700]; % mmol/L

% Filtrate into RC/Bowman's capsule for a healthy kidney
% A healthy kidney has a single-nephron GFR of approximately 79 +/- 42
nanoliters per minute (nL/min)
snGFR = 79 / 1000; % nL/min * 1e-3 = mL/min (filtrate into RC/Bowman's
capsule for a healthy kidney)

% Healthy
[C0_h, sn_h] = test_cases(C0, snGFR, 'healthy');
kidney_model(C0_h, sn_h, "Healthy");

% CKD3b
[C0_ckd, sn_ckd] = test_cases(C0, snGFR, 'ckd3b');
kidney_model(C0_ckd, sn_ckd, "CKD3b");

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% T2DM Early
[CO_dme, sn_dme] = test_cases(C0, snGFR, 't2dm_early');
kidney_model(C0_dme, sn_dme, "T2DM (Early)");

% T2DM Late
[CO_dml, sn_dml] = test_cases(C0, snGFR, 't2dm_late');
kidney_model(C0_dml, sn_dml, "T2DM (Late)");

% Hypertension
[CO_htn, sn_htn] = test_cases(C0, snGFR, 'htn');
kidney_model(C0_htn, sn_htn, "Hypertension");

% Combination of all 3, modeling severe CKD
[CO_tmp, sn_tmp] = test_cases(C0, snGFR, 'ckd3b');
[CO_tmp, sn_tmp] = test_cases(C0_tmp, sn_tmp, 'htn');
[CO_combo, sn_combo] = test_cases(C0_tmp, sn_tmp, 't2dm_late');
kidney_model(C0_combo, sn_combo, "CKD3b + HTN + T2DM (Late)");

```

#### Healthy

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##### 8x10 Molar Flow Rate Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> Columns 1 through 7

|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.0111 | 0.0081 | 0.0005 | 0.0004 | 0.0003 | 0.0019 | 0.0001 |
| 0.0074 | 0.0054 | 0.0005 | 0.0000 | 0.0003 | 0.0004 | 0.0001 |
| 0.0050 | 0.0036 | 0.0005 | 0.0000 | 0.0003 | 0.0003 | 0.0001 |
| 0.0039 | 0.0028 | 0.0002 | 0.0000 | 0.0001 | 0.0003 | 0.0001 |
| 0.0039 | 0.0028 | 0.0003 | 0.0000 | 0.0001 | 0.0003 | 0.0001 |
| 0.0029 | 0.0021 | 0.0003 | 0.0000 | 0.0001 | 0.0003 | 0.0000 |
| 0.0028 | 0.0020 | 0.0003 | 0.0000 | 0.0001 | 0.0003 | 0.0000 |
| 0.0028 | 0.0020 | 0.0003 | 0.0000 | 0.0001 | 0.0003 | 0.0000 |

##### Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |

##### 8x10 Concentrations Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> Columns 1 through 7

|          |          |        |        |        |         |        |
|----------|----------|--------|--------|--------|---------|--------|
| 140.0000 | 102.0000 | 5.7140 | 4.6905 | 4.3500 | 24.0000 | 0.8225 |
|----------|----------|--------|--------|--------|---------|--------|

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|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.0938 | 0.0683 | 0.0057 | 0.0005 | 0.0043 | 0.0048 | 0.0008 |
| 0.0630 | 0.0459 | 0.0057 | 0.0004 | 0.0043 | 0.0043 | 0.0008 |
| 0.0490 | 0.0357 | 0.0029 | 0.0004 | 0.0017 | 0.0043 | 0.0007 |
| 0.0490 | 0.0357 | 0.0033 | 0.0004 | 0.0017 | 0.0037 | 0.0007 |
| 0.0368 | 0.0268 | 0.0033 | 0.0004 | 0.0013 | 0.0037 | 0.0002 |
| 0.0349 | 0.0255 | 0.0033 | 0.0004 | 0.0014 | 0.0035 | 0.0002 |
| 0.0349 | 0.0255 | 0.0033 | 0.0004 | 0.0016 | 0.0035 | 0.0002 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.3950 | 0.0920 | 0.5700 |
| 0.0003 | 0.0001 | 0.0004 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |

Outlet (collecting duct) in grams/min per solute:

Na<sup>+</sup>: 6.3458e-05  
Cl<sup>-</sup>: 7.13e-05  
Urea: 1.5588e-05  
Glucose: 6.0749e-06  
K<sup>+</sup>: 4.8773e-06  
HCO<sub>3</sub><sup>-</sup>: 1.6816e-05  
Mg<sup>{2+}</sup>: 3.6008e-07  
PO<sub>4</sub><sup>{3-}</sup>: 1.2353e-06  
Creatinine: 1.0688e-06  
Ca<sup>{2+}</sup>: 5.4163e-07

CKD3b

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8x10 Molar Flow Rate Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup>

Columns 1 through 7

|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.0039 | 0.0028 | 0.0002 | 0.0001 | 0.0001 | 0.0006 | 0.0000 |
| 0.0026 | 0.0019 | 0.0002 | 0.0000 | 0.0001 | 0.0001 | 0.0000 |
| 0.0018 | 0.0013 | 0.0002 | 0.0000 | 0.0001 | 0.0001 | 0.0000 |
| 0.0014 | 0.0010 | 0.0001 | 0.0000 | 0.0001 | 0.0001 | 0.0000 |
| 0.0014 | 0.0010 | 0.0001 | 0.0000 | 0.0001 | 0.0001 | 0.0000 |
| 0.0010 | 0.0007 | 0.0001 | 0.0000 | 0.0000 | 0.0001 | 0.0000 |
| 0.0010 | 0.0007 | 0.0001 | 0.0000 | 0.0000 | 0.0001 | 0.0000 |
| 0.0010 | 0.0007 | 0.0001 | 0.0000 | 0.0001 | 0.0001 | 0.0000 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |

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|        |        |        |
|--------|--------|--------|
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |

#### 8x10 Concentrations Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na+, Cl-, Urea, Glucose, K+, HCO3-, Mg2+, PO4^3-, Creatinine, Ca2+ Columns 1 through 7

|          |          |        |        |        |         |        |
|----------|----------|--------|--------|--------|---------|--------|
| 140.0000 | 102.0000 | 7.9996 | 4.6905 | 5.0025 | 21.6000 | 0.8225 |
| 0.0938   | 0.0683   | 0.0080 | 0.0005 | 0.0050 | 0.0043  | 0.0008 |
| 0.0630   | 0.0459   | 0.0080 | 0.0004 | 0.0050 | 0.0039  | 0.0008 |
| 0.0490   | 0.0357   | 0.0040 | 0.0004 | 0.0020 | 0.0039  | 0.0007 |
| 0.0490   | 0.0357   | 0.0046 | 0.0004 | 0.0020 | 0.0033  | 0.0007 |
| 0.0368   | 0.0268   | 0.0046 | 0.0004 | 0.0015 | 0.0033  | 0.0002 |
| 0.0349   | 0.0255   | 0.0046 | 0.0004 | 0.0017 | 0.0031  | 0.0002 |
| 0.0349   | 0.0255   | 0.0046 | 0.0004 | 0.0018 | 0.0031  | 0.0002 |

#### Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.4938 | 0.1472 | 0.5130 |
| 0.0003 | 0.0001 | 0.0003 |
| 0.0002 | 0.0002 | 0.0003 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |

#### Outlet (collecting duct) in grams/min per solute:

Na<sup>+</sup>: 2.2361e-05  
 Cl<sup>-</sup>: 2.5125e-05  
 Urea: 7.6901e-06  
 Glucose: 2.1407e-06  
 K<sup>+</sup>: 1.9765e-06  
 HCO<sub>3</sub><sup>-</sup>: 5.3331e-06  
 Mg<sup>{2+}</sup>: 1.2688e-07  
 PO<sub>4</sub><sup>{3-}</sup>: 5.441e-07  
 Creatinine: 6.026e-07  
 Ca<sup>{2+}</sup>: 1.7177e-07

#### T2DM (Early)

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#### 8x10 Molar Flow Rate Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na+, Cl-, Urea, Glucose, K+, HCO3-, Mg2+, PO4^3-, Creatinine, Ca2+ Columns 1 through 7

|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.0124 | 0.0091 | 0.0005 | 0.0006 | 0.0004 | 0.0021 | 0.0001 |
| 0.0083 | 0.0061 | 0.0005 | 0.0001 | 0.0004 | 0.0004 | 0.0001 |
| 0.0056 | 0.0041 | 0.0005 | 0.0001 | 0.0004 | 0.0004 | 0.0001 |
| 0.0044 | 0.0032 | 0.0003 | 0.0001 | 0.0002 | 0.0004 | 0.0001 |

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|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.0044 | 0.0032 | 0.0003 | 0.0001 | 0.0002 | 0.0003 | 0.0001 |
| 0.0033 | 0.0024 | 0.0003 | 0.0001 | 0.0001 | 0.0003 | 0.0000 |
| 0.0031 | 0.0023 | 0.0003 | 0.0001 | 0.0001 | 0.0003 | 0.0000 |
| 0.0031 | 0.0023 | 0.0003 | 0.0001 | 0.0001 | 0.0003 | 0.0000 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.0000 | 0.0000 | 0.0001 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |

8x10 Concentrations Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> Columns 1 through 7

|          |          |        |        |        |         |        |
|----------|----------|--------|--------|--------|---------|--------|
| 140.0000 | 102.0000 | 5.7140 | 7.0000 | 4.3500 | 24.0000 | 0.8225 |
| 0.0938   | 0.0683   | 0.0057 | 0.0007 | 0.0043 | 0.0048  | 0.0008 |
| 0.0630   | 0.0459   | 0.0057 | 0.0006 | 0.0043 | 0.0043  | 0.0008 |
| 0.0490   | 0.0357   | 0.0029 | 0.0006 | 0.0017 | 0.0043  | 0.0007 |
| 0.0490   | 0.0357   | 0.0033 | 0.0006 | 0.0017 | 0.0037  | 0.0007 |
| 0.0368   | 0.0268   | 0.0033 | 0.0006 | 0.0013 | 0.0037  | 0.0002 |
| 0.0349   | 0.0255   | 0.0033 | 0.0006 | 0.0014 | 0.0035  | 0.0002 |
| 0.0349   | 0.0255   | 0.0033 | 0.0006 | 0.0016 | 0.0035  | 0.0002 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.3950 | 0.0920 | 0.5700 |
| 0.0003 | 0.0001 | 0.0004 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |

Outlet (collecting duct) in grams/min per solute:

Na<sup>+</sup>: 7.139e-05  
Cl<sup>-</sup>: 8.0212e-05  
Urea: 1.7537e-05  
Glucose: 1.0199e-05  
K<sup>+</sup>: 5.4869e-06  
HCO<sub>3</sub><sup>-</sup>: 1.8918e-05  
Mg<sup>{2+}</sup>: 4.0508e-07  
PO<sub>4</sub><sup>{3-}</sup>: 1.3897e-06  
Creatinine: 1.2024e-06  
Ca<sup>{2+}</sup>: 6.0933e-07

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T2DM (Late)

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8x10 Molar Flow Rate Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> Columns 1 through 7

|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.0023 | 0.0017 | 0.0001 | 0.0002 | 0.0001 | 0.0003 | 0.0000 |
| 0.0016 | 0.0011 | 0.0001 | 0.0000 | 0.0001 | 0.0001 | 0.0000 |
| 0.0010 | 0.0008 | 0.0001 | 0.0000 | 0.0001 | 0.0001 | 0.0000 |
| 0.0008 | 0.0006 | 0.0000 | 0.0000 | 0.0000 | 0.0001 | 0.0000 |
| 0.0008 | 0.0006 | 0.0001 | 0.0000 | 0.0000 | 0.0001 | 0.0000 |
| 0.0006 | 0.0004 | 0.0001 | 0.0000 | 0.0000 | 0.0001 | 0.0000 |
| 0.0006 | 0.0004 | 0.0001 | 0.0000 | 0.0000 | 0.0000 | 0.0000 |
| 0.0006 | 0.0004 | 0.0001 | 0.0000 | 0.0000 | 0.0000 | 0.0000 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |

8x10 Concentrations Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> Columns 1 through 7

|          |          |        |        |        |         |        |
|----------|----------|--------|--------|--------|---------|--------|
| 140.0000 | 102.0000 | 5.7140 | 9.7000 | 5.6550 | 20.4000 | 0.8225 |
| 0.0938   | 0.0683   | 0.0057 | 0.0010 | 0.0057 | 0.0041  | 0.0008 |
| 0.0630   | 0.0459   | 0.0057 | 0.0009 | 0.0057 | 0.0037  | 0.0008 |
| 0.0490   | 0.0357   | 0.0029 | 0.0009 | 0.0023 | 0.0037  | 0.0007 |
| 0.0490   | 0.0357   | 0.0033 | 0.0009 | 0.0023 | 0.0031  | 0.0007 |
| 0.0368   | 0.0268   | 0.0033 | 0.0009 | 0.0017 | 0.0031  | 0.0002 |
| 0.0349   | 0.0255   | 0.0033 | 0.0009 | 0.0019 | 0.0030  | 0.0002 |
| 0.0349   | 0.0255   | 0.0033 | 0.0009 | 0.0021 | 0.0030  | 0.0002 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.4542 | 0.0920 | 0.5700 |
| 0.0003 | 0.0001 | 0.0004 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |

Outlet (collecting duct) in grams/min per solute:

Na<sup>+</sup>: 1.3296e-05

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Cl<sup>-</sup>: 1.4939e-05  
Urea: 3.2661e-06  
Glucose: 2.6322e-06  
K<sup>+</sup>: 1.3285e-06  
HCO<sub>3</sub><sup>-</sup>: 2.9949e-06  
Mg<sup>{2+}</sup>: 7.5444e-08  
PO<sub>4</sub><sup>{3-}</sup>: 2.9764e-07  
Creatinine: 2.2394e-07  
Ca<sup>{2+}</sup>: 1.1348e-07

#### Hypertension

##### 8x10 Molar Flow Rate Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> Columns 1 through 7

|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.0100 | 0.0073 | 0.0004 | 0.0003 | 0.0003 | 0.0017 | 0.0001 |
| 0.0067 | 0.0049 | 0.0004 | 0.0000 | 0.0003 | 0.0003 | 0.0001 |
| 0.0045 | 0.0033 | 0.0004 | 0.0000 | 0.0003 | 0.0003 | 0.0001 |
| 0.0035 | 0.0025 | 0.0002 | 0.0000 | 0.0001 | 0.0003 | 0.0000 |
| 0.0035 | 0.0025 | 0.0002 | 0.0000 | 0.0001 | 0.0003 | 0.0000 |
| 0.0026 | 0.0019 | 0.0002 | 0.0000 | 0.0001 | 0.0003 | 0.0000 |
| 0.0025 | 0.0018 | 0.0002 | 0.0000 | 0.0001 | 0.0002 | 0.0000 |
| 0.0025 | 0.0018 | 0.0002 | 0.0000 | 0.0001 | 0.0002 | 0.0000 |

##### Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |
| 0.0000 | 0.0000 | 0.0000 |

##### 8x10 Concentrations Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> Columns 1 through 7

|          |          |        |        |        |         |        |
|----------|----------|--------|--------|--------|---------|--------|
| 140.0000 | 102.0000 | 5.7140 | 4.6905 | 4.3500 | 24.0000 | 0.8225 |
| 0.0938   | 0.0683   | 0.0057 | 0.0005 | 0.0043 | 0.0048  | 0.0008 |
| 0.0630   | 0.0459   | 0.0057 | 0.0004 | 0.0043 | 0.0043  | 0.0008 |
| 0.0490   | 0.0357   | 0.0029 | 0.0004 | 0.0017 | 0.0043  | 0.0007 |
| 0.0490   | 0.0357   | 0.0033 | 0.0004 | 0.0017 | 0.0037  | 0.0007 |
| 0.0368   | 0.0268   | 0.0033 | 0.0004 | 0.0013 | 0.0037  | 0.0002 |
| 0.0349   | 0.0255   | 0.0033 | 0.0004 | 0.0014 | 0.0035  | 0.0002 |
| 0.0349   | 0.0255   | 0.0033 | 0.0004 | 0.0016 | 0.0035  | 0.0002 |

##### Columns 8 through 10

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|        |        |        |
|--------|--------|--------|
| 0.3950 | 0.0920 | 0.5700 |
| 0.0003 | 0.0001 | 0.0004 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0003 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |
| 0.0002 | 0.0001 | 0.0002 |

Outlet (collecting duct) in grams/min per solute:

Na<sup>+</sup>: 5.7112e-05  
 Cl<sup>-</sup>: 6.417e-05  
 Urea: 1.4029e-05  
 Glucose: 5.4674e-06  
 K<sup>+</sup>: 4.3896e-06  
 HCO<sub>3</sub><sup>-</sup>: 1.5135e-05  
 Mg<sup>{2+}</sup>: 3.2407e-07  
 PO<sub>4</sub><sup>{3-}</sup>: 1.1117e-06  
 Creatinine: 9.6192e-07  
 Ca<sup>{2+}</sup>: 4.8747e-07

CKD3b + HTN + T2DM (Late)

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8x10 Molar Flow Rate Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine, Ca<sup>2+</sup> 1.0e-03 \*

Columns 1 through 7

|        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|
| 0.7349 | 0.5354 | 0.0420 | 0.0509 | 0.0341 | 0.0964 | 0.0043 |
| 0.4924 | 0.3587 | 0.0420 | 0.0051 | 0.0341 | 0.0193 | 0.0043 |
| 0.3309 | 0.2411 | 0.0420 | 0.0046 | 0.0341 | 0.0173 | 0.0043 |
| 0.2574 | 0.1876 | 0.0210 | 0.0046 | 0.0137 | 0.0173 | 0.0035 |
| 0.2574 | 0.1876 | 0.0241 | 0.0046 | 0.0137 | 0.0147 | 0.0035 |
| 0.1931 | 0.1407 | 0.0241 | 0.0046 | 0.0102 | 0.0147 | 0.0010 |
| 0.1834 | 0.1336 | 0.0241 | 0.0046 | 0.0113 | 0.0140 | 0.0010 |
| 0.1834 | 0.1336 | 0.0241 | 0.0046 | 0.0124 | 0.0140 | 0.0010 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.0030 | 0.0008 | 0.0027 |
| 0.0019 | 0.0008 | 0.0018 |
| 0.0015 | 0.0010 | 0.0013 |
| 0.0013 | 0.0010 | 0.0012 |
| 0.0013 | 0.0010 | 0.0012 |
| 0.0013 | 0.0010 | 0.0009 |
| 0.0012 | 0.0010 | 0.0009 |
| 0.0012 | 0.0010 | 0.0008 |

8x10 Concentrations Matrix

- Rows = RC, S1, S2, S3, DL, AL, DT, CD
- Columns = Na<sup>+</sup>, Cl<sup>-</sup>, Urea, Glucose, K<sup>+</sup>, HCO<sub>3</sub><sup>-</sup>, Mg<sup>2+</sup>, PO<sub>4</sub><sup>3-</sup>, Creatinine,

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Ca2+ Columns 1 through 7

|          |          |        |        |        |         |        |
|----------|----------|--------|--------|--------|---------|--------|
| 140.0000 | 102.0000 | 7.9996 | 9.7000 | 6.5032 | 18.3600 | 0.8225 |
| 0.0938   | 0.0683   | 0.0080 | 0.0010 | 0.0065 | 0.0037  | 0.0008 |
| 0.0630   | 0.0459   | 0.0080 | 0.0009 | 0.0065 | 0.0033  | 0.0008 |
| 0.0490   | 0.0357   | 0.0040 | 0.0009 | 0.0026 | 0.0033  | 0.0007 |
| 0.0490   | 0.0357   | 0.0046 | 0.0009 | 0.0026 | 0.0028  | 0.0007 |
| 0.0368   | 0.0268   | 0.0046 | 0.0009 | 0.0020 | 0.0028  | 0.0002 |
| 0.0349   | 0.0255   | 0.0046 | 0.0009 | 0.0021 | 0.0027  | 0.0002 |
| 0.0349   | 0.0255   | 0.0046 | 0.0009 | 0.0024 | 0.0027  | 0.0002 |

Columns 8 through 10

|        |        |        |
|--------|--------|--------|
| 0.5678 | 0.1472 | 0.5130 |
| 0.0004 | 0.0001 | 0.0003 |
| 0.0003 | 0.0002 | 0.0003 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |
| 0.0002 | 0.0002 | 0.0002 |

Outlet (collecting duct) in grams/min per solute:

Na<sup>+</sup>: 4.2167e-06

Cl<sup>-</sup>: 4.7378e-06

Urea: 1.4501e-06

Glucose: 8.3479e-07

K<sup>+</sup>: 4.8451e-07

HCO<sub>3</sub><sup>-</sup>: 8.5482e-07

Mg<sup>{2+}</sup>: 2.3927e-08

PO<sub>4</sub><sup>{3-}</sup>: 1.1799e-07

Creatinine: 1.1363e-07

Ca<sup>{2+}</sup>: 3.2392e-08







